### Figure 1A

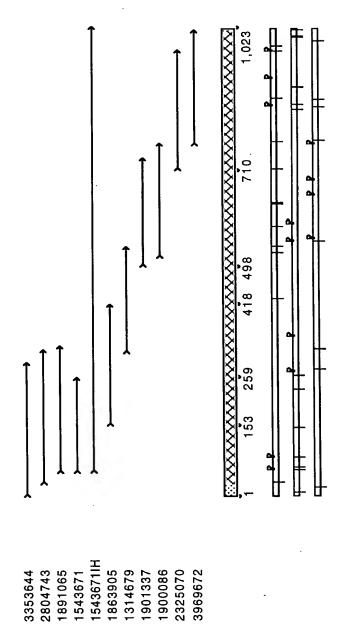
>3353644	CATTTGAGGC	CATATAAAGT	CACCTGAGGC	NCTCTCCACC	ACAGCCCACC
>2804743			GAGGC	CCTCTCCACC	ACAGCCCACC
>1891065					С
Consensus	CATTTGAGGC	CATATAAAGT	CACCTGAGGC	CCTCTCCACC	ACAGCCCACC
>3353644	AGTGACCATG	AAGGCTGTGC	TGCTTGCCCT	GTTGATGGCA	GGCŤTGGCCC
>2804743	AGTGACCATG	AAGGCTGTGC	TGCTTGCCCT	GTTGATGGCA	GGCTTGGCCC
>1891065	AGTGACCATG	AAGGCTGTGC	TGCTTGCCCT	GTTGATGGCA	GGCTTGGCCC
>1543671	GTGACCATG	AAGGCTGTGC	TGCTTGCCCT	GTTGATGGCA	GGCTTGGCCC
>1543671IH	GTGACCATG	AAGGCTGTGC	TGCTTGCCCT	GTTGATGGCA	GGCTTGGCCC
Consensus	AGTGACCATG	AAGGCTGTGC	TGCTTGCCCT	GTTGATGGCA	GGCTTGGCCC
>3353644	TGCAGCCAGG	CACTGCCCTG	CTGTGCTACT	CCTGCAAAGC	CCAGGTGAGC
>2804743	TGCAGCCAGG	CACTGCCCTG	CTGTGCTACT	CCTGCAAAGC	CCAGGTGAGC
>1891065	TGCAGCCAGG	CACTGCCCTG		CCTGCAAAGC	
>1543671	TGCAGCCAGG	CACTGCCCTG	CTGTGCTACT	CCTGCAAAGC	CCAGGTGAGC
>1543671IH	TGCAGCCAGG	CACTGCCCTG	CTGTGCTACT	CCTGCAAAGC	CCAGGTGAGC
Consensus	TGCAGCCAGG	CACTGCCCTG	CTGTGCTACT	CCTGCAAAGC	CCAGGTGAGC
. 2252644	* * CC * CC * CT	GCCTGCAGGT	CCACAACTICC	ACCCAGCTGG	CCCACCACTC
>3353644		GCCTGCAGGT		ACCCAGCTGG	
>2804743		GCCTGCAGGT		ACCCAGCTGG	
>1891065			001101010	ACCCAGCTGG	
>1543671		GCCTGCAGGT		ACCCAGCTGG	
>1543671IH		GCCTGCAGGT			
>1863905	• • • • • • • • • • • • • • • • • • • •	GCCTGCAGGT		ACCCAGCTGG	
Consensus	AACGAGGACT	GCCTGCAGGT	GGAGAACTGC	ACCCAGCTGG	GGGAGCAGTG
>3353644	CTGGACCGCG	CGCATCCGCG	CAGTTGGCCT	CNTGACCGTC	ATCAGCAAAG
>2804743	CTGGACCGCG	CGCATCCGCG	CAGTTGGCCT	CCTGACCGTC	ATCAGCAAAG
>1891065	CTGGACCGCG	CGCATCCGCG	CAGTTGGCCT	CCTGACCGTC	ATCAGCAAAG
>1543671	CTGGACCGCG	CGCATCCGCG	CAGTTGGCCT	CCTGACCGTC	ATCAGCAAAG
>1543671IH	CTGGACCGCG	CGCATCCGCG	CAGTTGGCCT	CCTGACCGTC	ATCAGCAAAG
>1863905	CTGGACCGCG	CGCATCCGCG	CAGTTGGCCT	CCTGACCGTC	ATCAGCAAAG
Consensus	CTGGACCGCG	CGCATCCGCG	CAGTTGGCCT	CCTGACCGTC	ATCAGCAAAG
2252644	omaa) aamm	0 omoodma	CA MCANIMOA C	AGGANTACTA	C
>3353644	: CTGCAGCTT		= '		
>2804743	GCTGCAGCTT			AGGACTACTA	
>1891065	GCTGCAGCTT	GAACTGCGTG	GATGACTCAC	AGGACTACTA	CGTGGGCAAG
>1543671	GCTGCAGC		a. ma. ama. a	1 CC 1 CM 1 CM 2	COMOCOON NO
>1543671IH					CGTGGGCAAG
>1863905					CGTGGGCAAG
Consensus	GCTGCAGCTT	GAACTGCGTG	GATGACTCAC	AGGACTACTA	CGTGGGCAAG
>2804743	AAGAACATCA	CGTGCTGT			
>1891065		CGTGCTGTGA	CACCGACT		
>1543671IH	AAGAACATCA	CGTGCTGTGA	CACCGACTTG	TGCAACGCCA	GCGGGGCCCA
>1863905	AAGAACATCA	CGTGCTGTGA	CACCGACTTG	TGCAACGCCA	GCGGGGCCCA
>1314679		GTGCTGTGA	CACCGACTTG	TGCAACGCCA	GCGGGGCCCA
Consensus	AAGAACATCA	CGTGCTGTGA	CACCGACTTG	TGCAACGCCA	GCGGGGCCCA

### Figure 1B

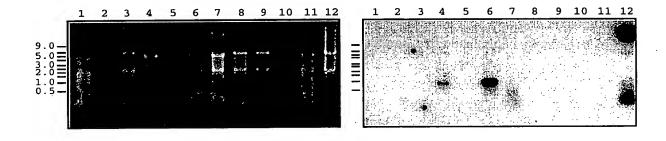
					,	
:	>1543671IH		+	CCATCCTTGC		
	>1863905	TGCCCTGCAG	CCGGCTGCCG	CCATCCTTGC	GCTGCTCCCT	GCACTCGGCC
:	>1314679	TGCCCTGCAG	CCGGCTGCCG	CCATCCTTGC	GCTGCTCCCT	GCACTCGGCC
	Consensus	TGCCCTGCAG	CCGGCTGCCG	CCATCCTTGC	GCTGCTCCCT	GCACTCGGCC
	>1543671IH	TGCTGCTCTG	GGGACCCGGC	CAGCTATAGG	CTCTGGGGGG	CCCCGCTGCA
	>1863905	TGCTGCTCTG				
	>1314679			CAGCTATAGG	CTCTGGGGGG	CCCCGCTGCA
	Consensus			CAGCTATAGG		
	Conscisus	1001001010				
	>1543671IH	GCCCACACTG	GGTGTGGTGC	CCCAGGCCTC	TGTGCCACTC	CTCACAGACC
	>1314679			CCCAGGCCTC		
	>1914075	0000	0010100100	••••		C
	Consensus	CCCCACACTG	ССТСТССТССТСС	CCCAGGCCTC	TGTGCCACTC	
	Consensus	GCCCACACIO	0010100100	0001.000010	101000	0.0
	>1543671IH	TCCCCCACTC	CCACCCTCTC	CTGGTTCCTG	AGGCACATCC	ТААСССААСТ
				CTGGTTCCTG		
	>1314679			CTGGTTCCTG		
	>1901337	TGGCCCAGTG		CTGGTTCCTG		
	>1900086			CTGGTTCCTG		
	Consensus	TGGCCCAGTG	GGAGCCTGTC	CTGGTTCCTG	AGGCACATCC	TAACGCAAGT
		cma1 aa1 mam	1 mamamaa 1 a	OCCUPATION OCC	CACCCTCACC	CTCCCATCCC
	>1543671IH			CCCTGTCCCC		
	>1901337			CCCTGTCCCC		
	>1900086			CCCTGTNCCC		
	Consensus	CTGACCATGT	ATGTCTGCAC	CCCTGTCCCC	CACCCTGACC	CTCCCATGGC
					omoma omoa o	1010100000
	>1543671IH			GGCAGATCAG		
	>1901337			GGCAGATCAG		
	>1900086			GGCAGATCAG		
	Consensus	CCTCTCCAGG	ACTCCCACCC	GGCAGATCAG	CTCTAGTGAC	ACAGATCCGC
					~~~~~~~~~~	maaaaaaa aa a
	>1543671IH			CCTCTCTGCT		
	>1901337					TGGCCCAGCA
	>1900086					TGGCCCAGCA
	Consensus	CTGCAGATGG	CCCCTCCAAC	CCTCTCTGCT	GCTGTTTCCA	TGGCCCAGCA
	•					1001100000
	>1543671IH			GCTCAGGCAC		AGGAAGCCTT
	>1901337			GCTCAGGCAC		
	>1900086	TTCTCCACCC				AGGAAGCCTT
	>2325070					AGGAAGCCTT
	Consensus	TTCTCCACCC	TTAACCCTGT	GCTCAGGCAC	CTCTTCCCCC	AGGAAGCCTT
	>1543671IH			ACTTGAGCCA	GGTCTGGTCC	GTGGTGTCCC
	>1900086	CCCTGCCCAC				
	>2325070	CCCTGCCCAC				GTGGTGTCCC
	>3969672					GTGGTGTCCC
	Consensus	CCCTGCCCAC	CCCATCTATG	ACTTGAGCCA	GGTCTGGTCC	GTGGTGTCCC
	>1543671IH					AAGGCTGAGA
	>2325070					AAGGCTGAGA
	>3969672					AAGGCTGAGA
	Consensus	CCGCACCCAG	CAGGGGACAG	GCACTCAGGA	GGGCCCAGTA	AAGGCTGAGA

# Figure 1C

>1543671IH	TGAAGTGGAC	TGAGTAGAAC	TGGAGGACAA	GAGTCGACGT	GAGTTCCTGG
>2325070	TGAAGTGGAC	TGAGTAGAAC	TGGAGGACAA	GAGTCGACGT	GAGTTCCTGG
>3969672	TGAAGTGGAC	TGAGTAGAAC	TGGAGGACAA	GAGTCGACGT	GAGTTCCTGG
Consensus	TGAAGTGGAC	TGAGTAGAAC	TGGAGGACAA	GAGTCGACGT	GAGTTCCTGG
>1543671IH	GAGTCTCCAG	AGATGGGGCC	TGGAGGCCTG	GAGGAAGGGG	CCAGGCCTCA
>2325070	GAGTCTCCAG	AGATGGGGCC	TGGAGGCCTG	GAGGAAGGGG	CCAGGCCTCA
>3969672	GAGTCTCCAG	AGATGGGGCC	TGGAGGCCTG	GAGGAAGGGG	CCAGGCCTCA
Consensus	GAGTCTCCAG	AGATGGGGCC	TGGAGGCCTG	GAGGAAGGGG	CCAGGCCTCA
>1543671IH	CATTCGTGGG	GCTCCCTGAA	TGGCAGCCTG	AGCACAGCGT	AGGCCCTTAA
>2325070	CATTCGTGGG	GCTCCCTGAA			
>3969672	CATTCGTGGG	GCTCCCTGAA	TGGCAGCCTG	AGCACAGCGT	AGGCCCTTAA
Consensus	CATTCGTGGG	GCTCCCTGAA	TGGCAGCCTG	AGCACAGCGT	AGGCCCTTAA
>1543671IH	TAAACACCTG	TTGGATAAGC	CCA		
>3969672	TAAACACCTG	TTGGAT			
Consensus	TAAACACCTG	TTGGATAAGC	CCA		

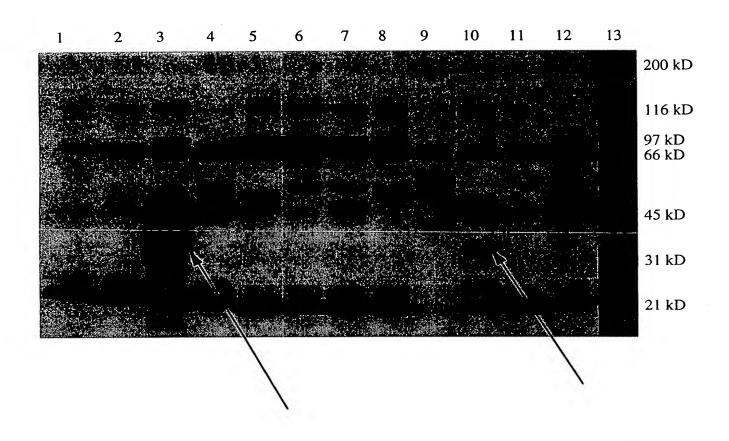


## Figure 3



Lane	Tissue	Lane	Tissue
1	Normal Prostate	7	Prostate Cancer
2	Normal Prostate	8	Prostate Cancer
3	Normal Prostate	9	Prostate Cancer
4	Normal Prostate	10	Prostate Cancer
5	Normal Prostate	11	Prostate Cancer
6	Normal Prostate	12	PS116 Bacterial RNA

### Figure 4



Lane	Tissue	Lane	Tissue
1	Bladder Cancer	8	Breast
2	Bladder	9	Prostate Cancer
3	Bladder Cancer	10	Prostate Cancer
4	Bladder	11	Prostsate Cancer
5	Bladder Cancer	12	ВРН
6	Colon	13	Markers
7	Lung		